REMARKS

The above amendments to the specification have been made to be in compliance with C.F.R. § 1.821-1.825. Accordingly, no new matter is added by these amendments.

Attached hereto is a marked-up version of the changes made to the specification and claims by the current amendment. The attached page is captioned "Version with markings to show changes made".

In the unlikely event that the fee transmittal is separated from this document and the Patent Office determines that an extension and/or other relief is required, Applicant petitions for any required relief including extensions of time and authorizes the Assistant Commissioner to charge the cost of such petitions and/or other fees due in connection with the filing of this document to **Deposit Account No. 03-1952** referencing docket no. 529872000100.

Respectfully submitted,

Dated: Syt 5, 2002

By:

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VERSION WITH MARKINGS TO SHOW CHANGES MADE

In the Specification:

Please substitute the following paragraphs.

On page 12, paragraph 50

[0051] Fig. 1 (SEQ ID NO:34) is a diagram showing the position of the *P[FRT]* insertion on the left arm of the *FRT* chromosome before being clipped according to a preferred embodiment of the present invention;

On page 13, paragraph 54, substitute the following paragraph.

[0055] Fig. 5 (SEQ ID NO:35) is a diagram showing the *P[FRT]* insertion on the right arm of the *FRT* chromosome before being clipped according to a preferred embodiment of the present invention; and

Please substitute the following table on page 18, with the following.

IR (1-20)	CATGATGAAATAACATAAGG (SEQ ID NO:1)
P5' sense (26-45)	CCGTCGAAAGCCGAAGCTTA (SEQ ID NO:2)
P5' antisens (264-232)	CCCAAGGCTCTGCTCCCACAATT (SEQ ID NO:3)
P3' sense (2756-2775)	AAACCCCACGGACATGCTAA (SEQ ID NO:4)
P3' (2878-2839)	CGGCAAGAGACATCCACTTA (SEQ ID NO:5)

Please substitute the following table on page 19, with the following.

Plac1	CACCCAAGGCTCTGCTCCCACAAT (SEQ ID NO:6)
Pryl	CCTTAGCATGTCCGTGGGGTTTGAAT (SEQ ID NO:7)
Pry2	CTTGCCGACGGGACCACCTTATGTTATT (SEQ ID NO:8)
Splac2	GAATTCACTGGCCGTCGTTTTACAA (SEQ ID NO:9)
Sp1	ACACAACCTTTCCTCTCAACAA (SEQ ID NO:10)
Sp3	GAGTACGCAAAGCTTTAACTATGT (SEQ ID NO:11)